-3-

Atkinson et al. Application No. 09/554,941

Pursuant to the Examiner's recommendation, Applicant has amended the specification and claims to indicate the SEQ ID NO's of each amino acid sequence. Applicant therefore believes that the instant application is now in full compliance with the requirements of 37 CFR 1.821-1.825, obviating the need for another substitute sequence listing.

Attached hereto is a marked-up version of the changes made to the specification and claims by the current amendment. The attached page is captioned "Version With Markings To Show Changes Made."

Applicant respectfully requests entry of this amendment and favorable consideration of the claims. If in the opinion of the Examiner, a telephone conference would expedite the prosecution of the subject application, the Examiner is invited to call the undersigned attorney.

Respectfully submitted,

J. Timothy Meigs

Attorney for Applicants Registration No. 38,241

Syngenta Biotechnology, Inc. Patent Department P. O. Box 12257

Research Triangle Park, NC 27709-2257

Telephone: 919-541-8587

Date: October 12, 2001

-4-

Atkinson et al. Application No. 09/554,941

## Version With Markings To Show Changes Made

## In the specification:

The bullet point at page 3, lines 9-10, has been replaced with the following:

the linker peptide is characterized by comprising the amino acid sequence
QASSYTAPQPQ (SEQ ID NO:2) --

The bullet point at page 3, lines 11-12, has been replaced with the following:

the linker peptide is characterized by comprising the amino acid sequence
VILGVGPAKIQFEG (SEQ ID NO:1) --

The bullet point at page 3, lines 13-14, has been replaced with the following:

the linker peptide is characterized by comprising the amino acid sequence
QASIEGRYTAPQPQ (SEQ ID NO:11) --

## In the claims:

Claims 9-11 have been rewritten as follows:

- 9. (Amended) The method according to claim 1, wherein the linker peptide is characterized by comprising the amino acid sequence QASSYTAPQPQ (SEQ ID NO:2).
- 10. (Amended) The method according to claim [7] 1, wherein the linker peptide is characterized by comprising the amino acid sequence VILGVGPAKIQFEG (SEQ ID NO:1).
- 11. (Amended) The method according to claim [7] 1, wherein the linker peptide is characterized by comprising the amino acid sequence QASIEGRYTAPQPQ (SEQ ID NO:11).